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<110> Sims. John E.
   Born. Teresa L.
   Smith. Dirk E.
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<151> 1998-12-14

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Gly Asn Leu Ile Ala Val Pro Asp Lys Asn Tyr Ile Arg Pro Glu Ile 50 55 60

Phe Phe Ala Leu Ala Ser Ser Leu Ser Ser Ala Ser Ala Glu Lys Gly 65 70 75 80

Ser Pro Ile Leu Leu Gly Val Ser Lys Gly Glu Phe Cys Leu Tyr Cys 85 90 95

Asp Lys Asp Lys Gly Gln Ser His Pro Ser Leu Gln Leu Lys Lys Glu 100 105 110

Lys Leu Met Lys Leu Ala Ala Gln Lys Glu Ser Ala Arg Arg Pro Phe 115 120 125

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Ser Leu Ala Gln Ser Ala Gly Leu Ser Leu Met Trp Tyr Lys Ser Ser 65 70 . 75 80

Ser Lys Glu Glu Asp Ser Ile Trp Phe Arg Pro Thr Leu Leu Gln Asp 100 105 110

Ser Gly Leu Tyr Ala Cys Val Ile Arg Asn Ser Thr Tyr Cys Met Lys
115 120 125

Val Ser Ile Ser Leu Thr Val Gly Glu Asn Asp Thr Gly Leu Cys Tyr 130 135 140

Asn Ser Lys Met Lys Tyr Phe Glu Lys Ala Glu Leu Ser Lys Ser Lys 145 150 155 160

Glu Ile Ser Cys Arg Asp Ile Glu Asp Phe Leu Leu Pro Thr Arg Glu
165 170 175

Pro Glu Ile Leu Trp Tyr Lys Glu Cys Arg Thr Lys Thr Trp Arg Pro 180 185 190

Ser Ile Val Phe Lys Arg Asp Thr Leu Leu Ile Arg Glu Val Arg Glu
195 200 205

Asp Asp Ile Gly Asn Tyr Thr Cys Glu Leu Lys Tyr Gly Gly Phe Val 210 215 220

225 Pro Thr Tyr Phe Ile 305	Arg Pro Gln Ser Ile 290 Leu	Lys Leu Gly 275 Glu	Leu Gly 260 Asp	Leu 245 Asp	230 Tyr Ser	Pro	Met	Glu	Ser 250	235 Lys	Leu	Thr		G1n 255	240
Pro Thr Tyr Phe 11e 305	Gln Ser Ile 290	Leu Gly 275 Glu	Gly 260 Asp	245 Asp	Tyr Ser				250	Lys			lle	255	
Thr Tyr Phe Ile 305	Gln Ser Ile 290	Leu Gly 275 Glu	Gly 260 Asp	245 Asp	Ser				250				He	255	Glu
Tyr Phe Ile 305	Ser Ile 290	Gly 275 Glu	260 Asp	Asp		Ala	Asn	Leu		•					
Tyr Phe Ile 305	Ser Ile 290	Gly 275 Glu	260 Asp			Ala	Asn	Leu	Thr		_	_			
Phe Ile 305	Ile 290	275 G1u	Asp	Val	Ser					Cys	Arg	Ala		Phe	Gly
Phe Ile 305	Ile 290	275 G1u		Val	Ser			265					270		
Ile 305	290	Glu	Asp			Pro	Leu	He	Tyr	Trp	Met	Lys	Gly	Glu	Lys
Ile 305	290		Asp				280					285			
305		Lys		Leu	Asp	Glu	Asn	Arg	Va 1	Trp	Glu	Ser	Asp	He	Arg
305	Leu	Lys				295					300				
			Glu	His	Leu	Gly	Glu	Gln	Glu	Val	Ser	He	Ser	Leu	Пe
Val					310					315					320
	Asp	Ser	Val	Glu	Glu	Gly	Asp	Leu	Gly	Asn	Tyr	Ser	Cys	Tyr	Val
				325					330					335	
Glu	Asn	Gly	Asn	Gly	Arg	Arg	His	Ala	Ser	Val	Leu	Leu	His	Lys	Arg
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Leu	Leu	Leu	Val	Cys	Leu	Val	Thr	He	Tyr	Lys	Cys	Tyr	Lys	He	Glu
	370					375					380				
He	Met	Leu	Phe	Tyr	Arg	Asn	His	Phe	Gly	Ala	Glu	Glu	Leu	Asp	Gly
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Asp	Asn	Lys	Asp	Tyr	Asp	Ala	Tyr	Leu	Ser	Tyr	Thr	Lys	Va1	Asp	Pro
				405					410					415	
Asp	Gln	Trp	Asn	GIn	Glu	Thr	Gly	Glu	Glu	Glu	Arg	Phe	Ala	Leu	Glu
			420					425					430		
He	Leu	Pro	Asp	Met	Leu	Glu	Lys	His	Tyr	Gly	Tyr	Lys	Leu	Phe	He
		435					440					445			
Pro	Asp	Arg	Asp	Leu	He	Pro	Thr	Gly	Thr	Tyr	Ile	Glu	Asp	Val	Ala
	450					455				•	460		•		
Arg	Cys	Val	Asp	Gln	Ser	Lys	Arq	Leu	He	He		Met	Thr	Pro	Asn
465			•		470	J	J		_	475					480
Tyr	Val	Val	Arg	Arq		Trp	Ser	He	Phe	_	l eu	Glu	Thr	Ara	
-	-	-	5	485	- · J		,	•	490	-				495	
Ara		Met	Fen		Thr	Glv	Glu	Πρ	-	Val	He	{ en	عاآ	_	Cvs
J	Asn	. •				- · J			-,, -	'		_ ~ ~	• • •		
Ser	Asn		500					505	·				510	~·u	-, 0

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His Thr Ile Lys Leu Leu Thr Val Ile Lys Trp His Gly Pro Lys Cys
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Asn Lys Leu Asn Ser Lys Phe Trp Lys Arg Leu Gln Tyr Glu Met Pro
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Glu Gln Gly Pro Phe Gly Glu Leu Gln Thr Val Ser Ala Ile Ser Met
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Ala Ala Ala Thr Ser Thr Ala Leu Ala Thr Ala His Pro Asp Leu Arg
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Ser Thr Phe His Asn Thr Tyr His Ser Gln Met Arg Gln Lys His Tyr
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                                             620
Tyr Arg Ser Tyr Glu Tyr Asp Val Pro Pro Thr Gly Thr Leu Pro Leu
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Thr Ser Ile Gly Asn Gln His Thr Tyr Cys Asn Ile Pro Met Thr Leu
                645
                                     650
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Ile Asn Gly Gln Arg Pro Gln Thr Lys Ser Ser Arg Glu Gln Asn Pro
            660
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Asp Glu Ala His Thr Asn Ser Ala Ile Leu Pro Leu Leu Pro Arg Glu
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Pro Leu Glu Pro Gly Pro Ser Leu Pro Thr Met Asn Phe Val His Thr
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Ser Pro Lys Val Lys Asn Leu Asn Pro Lys Lys Phe Ser Ile His Asp
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Gln Asp His Lys Val Leu Val Leu Asp Ser Gly Asn Leu Ile Ala Val
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Pro Asp Lys Asn Tyr Ile Arg Pro Glu Ile Phe Phe Ala Leu Ala Ser
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                                     90
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Ala Gln Lys Glu Ser Ala Arg Arg Pro Phe Ile Phe Tyr Arg Ala Gln
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Ile Cys Thr Ser Cys Asn Cys Asn Glu Pro Val Gly Val Thr Asp Lys
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                                 185
                                                      190
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                                  25
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                              40
                                                  45
Leu Val Leu Asp Ser Gly Asn Leu Ile Ala Val Pro Asp Lys Asn Tyr
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Ile Arg Pro Glu Ile Phe Phe Ala Leu Ala Ser Ser Leu Ser Ser Ala
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Ser Ala Glu Lys Gly Ser Pro Ile Leu Leu Gly Val Ser Lys Gly Glu
                 85
Phe Cys Leu Tyr Cys Asp Lys Asp Lys Gly Gln Ser His Pro Ser Leu
            100
                                 105
                                                     110
Gln Leu Lys Lys Glu Lys Leu Met Lys Leu Ala Ala Gln Lys Glu Ser
        115
                             120
                                                 125
Ala Arg Arg Pro Phe Ile Phe Tyr Arg Ala Gln Val Gly Ser Trp Asn
    130
                        135
                                             140
Met Leu Glu Ser Ala Ala His Pro Gly Trp Phe Ile Cys Thr Ser Cys
145
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Asn Cys Asn Glu Pro Val Gly Val Thr Asp Lys Phe Glu Asn Arg Lys
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                                                         175
His Ile Glu Phe Ser Phe Gln Pro Val Cys Lys Ala Glu Met Ser Pro
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<211> 157

<212> PRT

<213> Homo sapiens

<211> 27

<212> PRT

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                                                      30
                                 25
Leu Ala Ser Ser Leu Ser Ser Ala Ser Ala Glu Lys Gly Ser Pro Ile
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                                                  45
Leu Leu Gly Val Ser Lys Gly Glu Phe Cys Leu Tyr Cys Asp Lys Asp
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                                             60
Lys Gly Gln Ser His Pro Ser Leu Gln Leu Lys Lys Glu Lys Leu Met
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Lys Leu Ala Ala Gln Lys Glu Ser Ala Arg Arg Pro Phe Ile Phe Tyr
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Arg Ala Gln Val Gly Ser Trp Asn Met Leu Glu Ser Ala Ala His Pro
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Gly Trp Phe Ile Cys Thr Ser Cys Asn Cys Asn Glu Pro Val Gly Val
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Thr Asp Lys Phe Glu Asn Arg Lys His Ile Glu Phe Ser Phe Gln Pro
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Val Cys Lys Ala Glu Met Ser Pro Ser Glu Val Ser Asp
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11 <213> Artificial Sequence <220> <223> Description of Artificial Sequence: leucine zipper polypeptide <400> 12 Pro Asp Val Ala Ser Leu Arg Gln Gln Val Glu Ala Leu Gln Gly Gln 10 1 5 15 Val Gln His Leu Gln Ala Ala Phe Ser Gln Tyr 20 25 <210> 13 <211> 33 <212> PRT <213> Artificial Sequence <220> <223> Description of Artificial Sequence: leucine zipper polypeptide <400> 13 Arg Met Lys Gln Ile Glu Asp Lys Ile Glu Glu Ile Leu Ser Lys Ile 1 10 15 Tyr His Ile Glu Asn Glu Ile Ala Arg Ile Lys Lys Leu Ile Gly Glu 20 25 30 Arg <210> 14 <211> 8 <212> PRT <213> Artificial Sequence <220> <223> Description of Artificial Sequence: polymorphic sequence from exon 2 of Tango 77

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<210> 15

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<220>

<223> Description of Artificial Sequence: polymorphic
 sequence from exon 2 of Tango 77

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Pro Ala Val Ser Pro Leu Glu Pro

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